



SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Stuiver, Maarten
 Ponstein, Anne
 Ohl, Stephan
 Goddijn, Oscar
 Simons, Lambertus
 Dekker, Bernardus
 Hoekstra, Sietske
 Tigelaar, Hendrik
 Elzinga, Nicolas
- (ii) TITLE OF INVENTION: Plasmids For Plant Transformation And Method For Using The Same
- (iii) NUMBER OF SEQUENCES: 19
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Zeneca Ag Products Inc.
 - (B) STREET: 1800 Concord Pike
 - (C) CITY: Wilmington
 - (D) STATE: DE
 - (E) COUNTRY: USA
 - (F) ZIP: 19850
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 09/469,812
 - (B) FILING DATE: 22-DEC-1999
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: EP 97201990.5
 - (B) FILING DATE: 30-JUN-1997
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: PCT/EP98/04171
 - (B) FILING DATE: 29-JUN-1998
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Hohenschutz, Liza D.
 - (B) REGISTRATION NUMBER: 33,712
 - (C) REFERENCE/DOCKET NUMBER: MOG 57688/UST
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (302) 886-1699
- (2) INFORMATION FOR SEQ ID NO: 1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 21 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA

(iii)	ANTI-SENSE: NO		
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 1:		
GATGGCAA	GATGGCAAAC GCTAATAAGG G		
(2) INFO	RMATION FOR SEQ ID NO: 2:		
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear		
(ii)	MOLECULE TYPE: cDNA		
(iii)	HYPOTHETICAL: NO		
(iii)	ANTI-SENSE: NO		
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 2:		
GAGCGGGT'	TT AACCTACTTC C	21	
(2) INFO	RMATION FOR SEQ ID NO: 3:		
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 39 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear		
(ii)	MOLECULE TYPE: cDNA		
(iii)	HYPOTHETICAL: NO		
(iii)	ANTI-SENSE: NO		
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 3:		
CTGTTGTC	GA CGGCTGGCTG GTGGCAGGAT ATATTGTGG	39	
(2) INFO	RMATION FOR SEQ ID NO: 4:		
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 38 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear		
(ii)	MOLECULE TYPE: cDNA		
(iii)	HYPOTHETICAL: NO		
(iii)	ANTI-SENSE: NO		

(iii) HYPOTHETICAL: NO

(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 4:	
CCAGCCGTC	CG ACAACAGCTC CCCGACCGGC AGCTCGGC	38
(2) INFOR	RMATION FOR SEQ ID NO: 5:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 44 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(iii)	ANTI-SENSE: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 5:	
TCGACGGGG	T GCCTGGGGCG GTGGCGGGGC GTGC	44
(2) INFOR	MATION FOR SEQ ID NO: 6:	
(i) :	SEQUENCE CHARACTERISTICS: (A) LENGTH: 44 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) l	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(iii) <i>i</i>	ANTI-SENSE: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 6:	
TCGAGCACG	C CCCGGCCCGC CCCCAGGCAC CCCG	44
(2) INFOR	MATION FOR SEQ ID NO: 7:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: double (D) TOPOLOGY: linear	
(ii) 1	MOLECULE TYPE: cDNA	
(iii) H	HYPOTHETICAL: NO	
(iii) A	ANTI-SENSE: NO	
(xi) S	SEQUENCE DESCRIPTION: SEQ ID NO: 7:	
TTCTCGAGAC	G GGTCGGTTAC TCG	23
(2) INFORM	MATION FOR SEQ ID NO: 8:	

(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 23 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ'ID NO: 8:	
TTGTCGAC	TC CAAGACGCCG AGC	23
(2) INFO	RMATION FOR SEQ ID NO: 9:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 9:	
AATTCAGA:	TC TCCATGGATC GATGAGCT	28
(2) INFO	RMATION FOR SEQ ID NO: 10:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	
(iii)	HYPOTHETICAL: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 10:	
CATCGATC	CA TGGAGATCTG	20
(2) INFO	RMATION FOR SEQ ID NO: 11:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: cDNA	

(iii) HYPOTHETICAL: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:	
GATCGGTCGA CC	12
(2) INFORMATION FOR SEQ ID NO: 12:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 23 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(iii) HYPOTHETICAL: NO	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:	
GACGCTAAAG GCAAACTTGA TTC	23
(2) INFORMATION FOR SEQ ID NO: 13:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 23 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(iii) HYPOTHETICAL: NO	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:	
TTAACCTACT TCCTTTGGTT CCG	23
(2) INFORMATION FOR SEQ ID NO: 14:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 23 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(iii) HYPOTHETICAL: NO	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:	
AGCGGCCCAT TTTAGTAGAC GGG	23
(2) INFORMATION FOR SEQ ID NO: 15:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 28 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single	

(ii) MOLECULE TYPE: cDNA	
(iii) HYPOTHETICAL: NO	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:	
ATCAATTCGA ACCCCAGAGT CCCGTCCA	28
(2) INFORMATION FOR SEQ ID NO: 16:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 21 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(iii) HYPOTHETICAL: NO	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:	
CTGGATCGTT TCGCATGATT G	21
(2) INFORMATION FOR SEQ ID NO: 17:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 27 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(iii) HYPOTHETICAL: NO	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:	
CCAACTTATC AGTGATAAAG AATCCGC	27
(2) INFORMATION FOR SEQ ID NO: 18:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 27 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(iii) HYPOTHETICAL: NO	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:	

(D) TOPOLOGY: linear

GAGATCAGAT TGTCGTTTCC CGCCTTC

(2) INFORMATION FOR SEQ ID NO:19:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 12 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

TNCAATTGAA AY

12